

Fig. 1

A

Amin Acid Number: 770 782

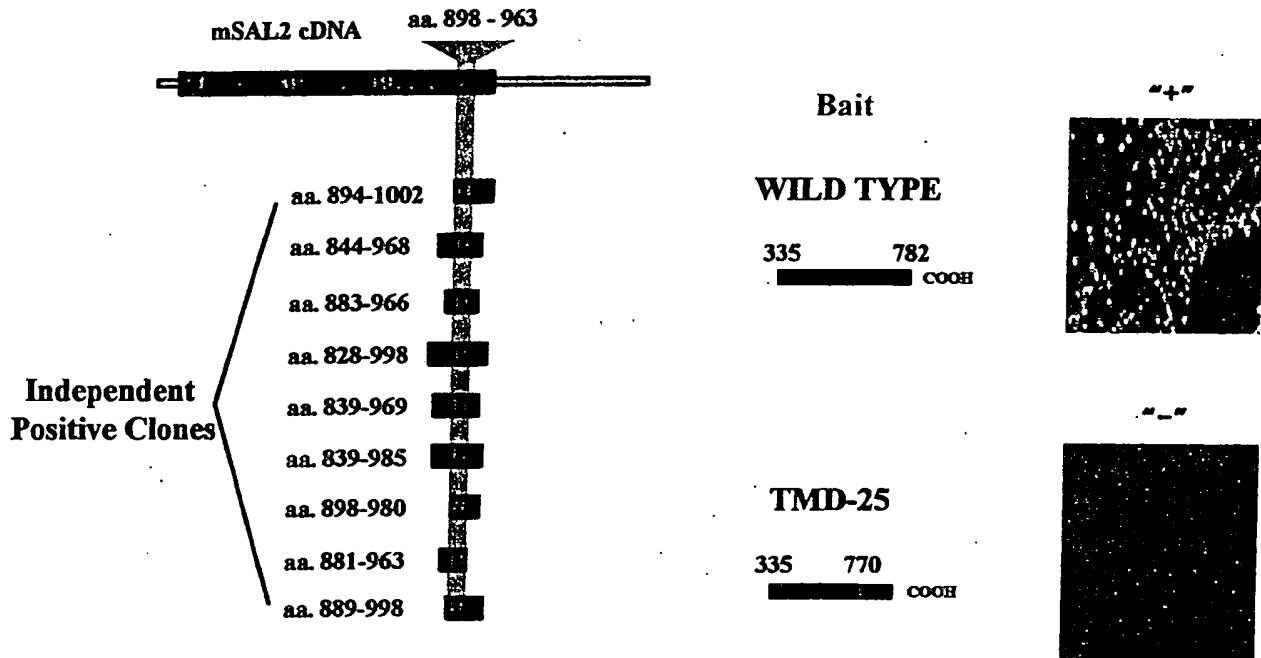
WT LT: - GAT ATA CTT TGT AAT GTG CAA GAA GGC GAC GAC CCC TTG AAG GAC ATA TGT GAA TAT AGC TGA

- D I L C N V Q E G D D P L R D I C E Y S *

- D I L C N V Q E G D F V M C R K A T T P *

TMD25 LT: - GAT ATA CTT TGT AAT GTG CAA GAA GAC TTT GTA ATG TGC AAG AAG GCG ACG ACC CCT TGA

B.



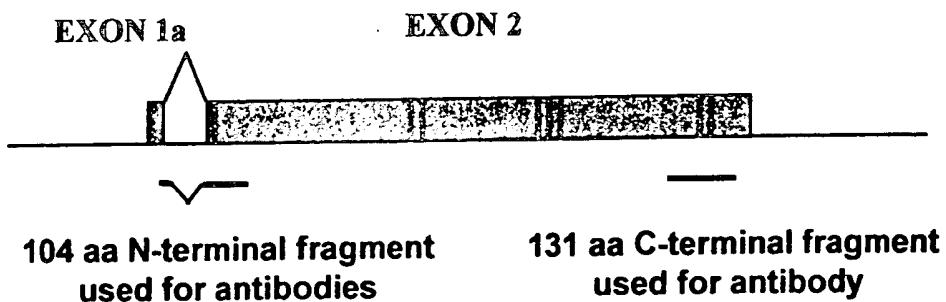
C.

Large T Deletions

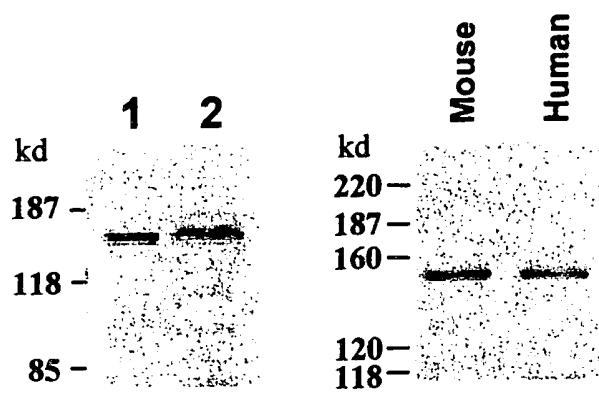
Wild Type	- N V Q E G D D P L R D I C E Y S *	+
335-780	- N V Q E G D D P L R D I C E * +	
335-776	- N V Q E G D D P L R * +	
335-774	- N V Q E G D D * -	-
335-770	- N V Q E * -	-
Δ 774	- N V Q E G D D - L R D I C E Y S * +	
Δ 775	- N V Q E G D D P - R D I C E Y S * +	
Δ 776	- N V Q E G D D P L - D I C E Y S * +	
Δ 774-776	- N V Q E G D D - - D I C E Y S * -	-

Fig. 2

A.



B.



C.

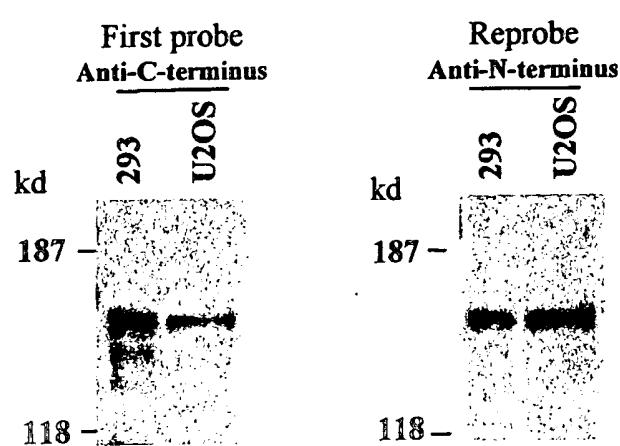


Fig. 3

A.

a b c d e f g h



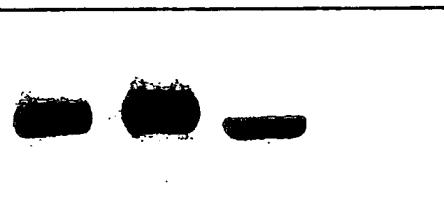
* Nonspecific
Wt LT
TMDLT

B.

Co-transfection

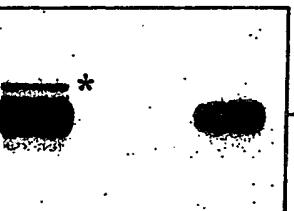
GST-mSAL2		GST-mSAL2	
/WT-LT	/TMD-LT	input	pulldown
input	pulldown	input	pulldown

Wt LT
TMDLT



Infection

Input	Normal IgG	Anti-mSAL2



GST-mSal2

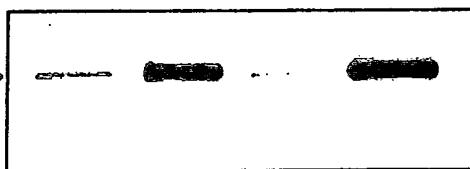


Fig. 4

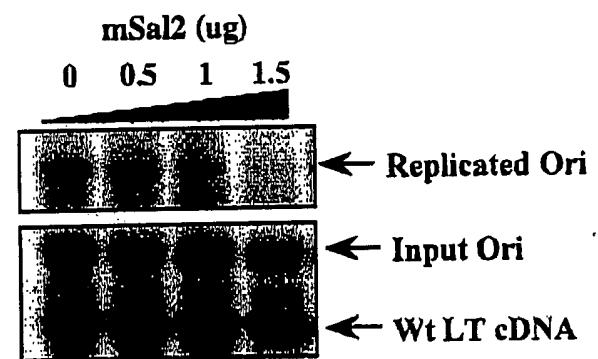
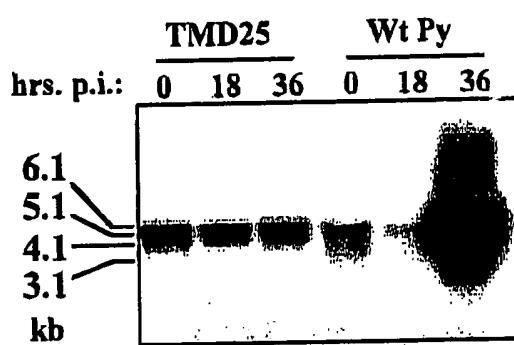
A



TMD-25

WILD TYPE

B



C

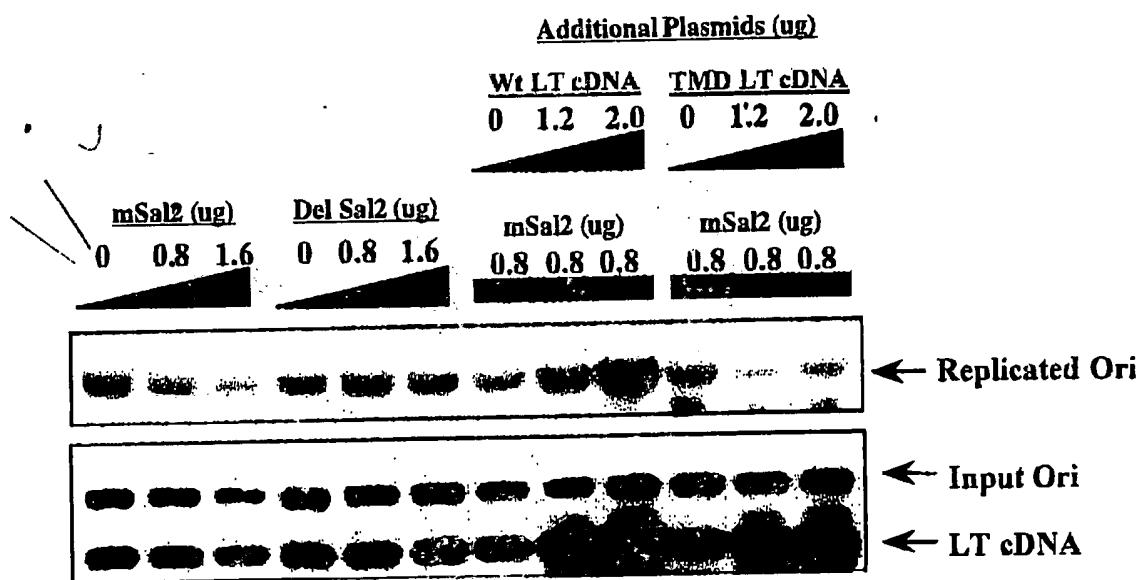


Figure 5

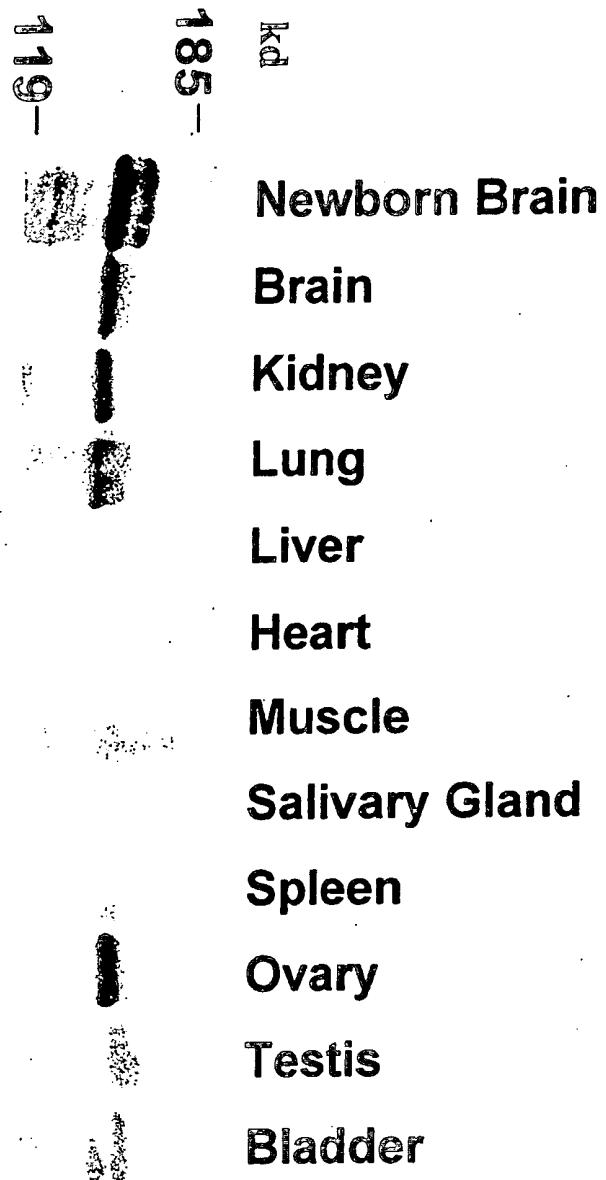
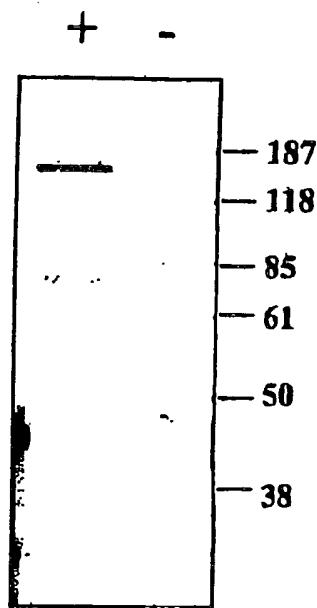


Fig. 6

N
327A
339A
404A
425A
429A
430A
472A
433B
420C
480
351A
352B
386A
504A
432A
517
520C
522A
523A
528A

Fig. 7



293 Cell extract

Fig. 8

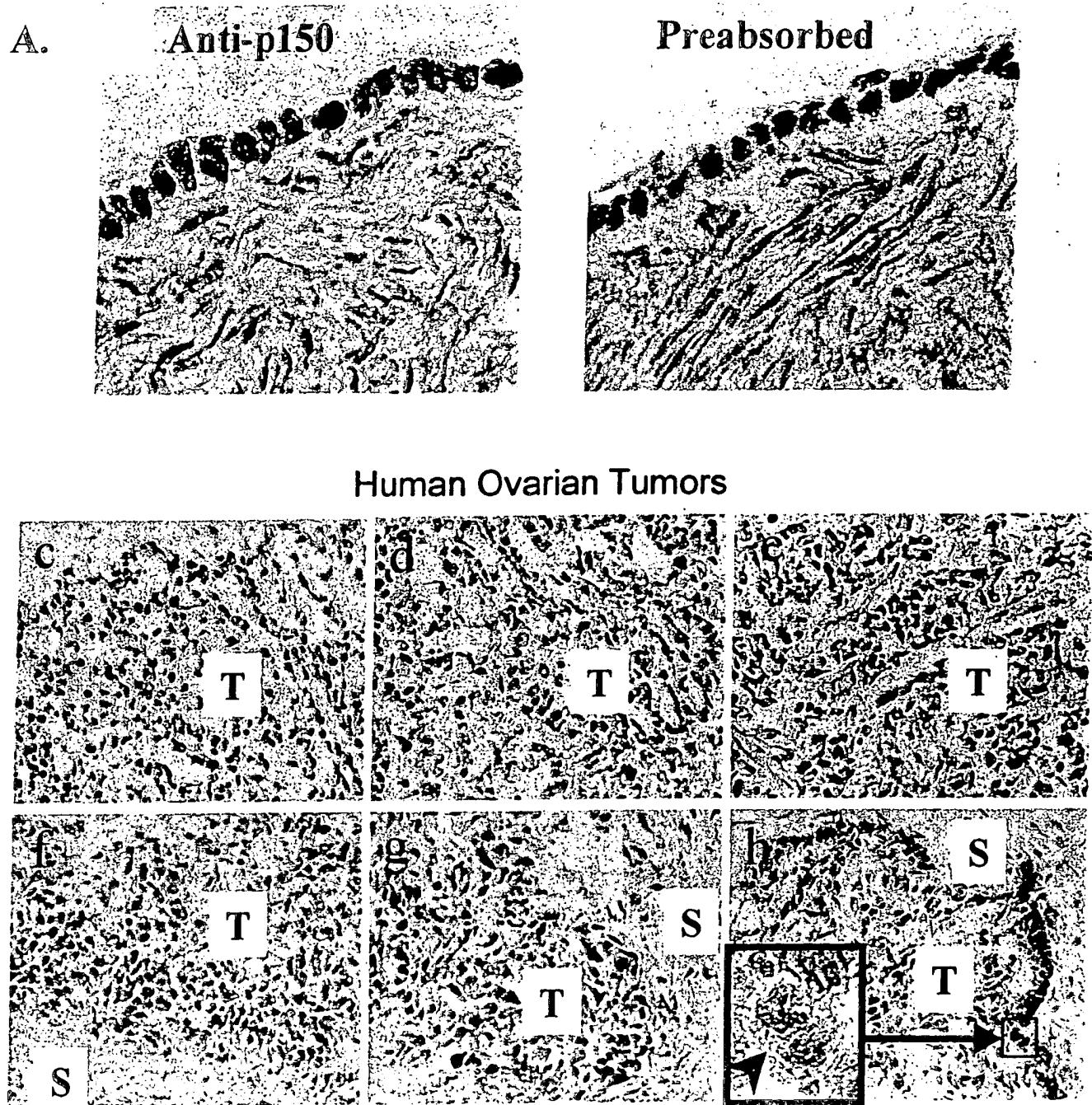


Fig. 9

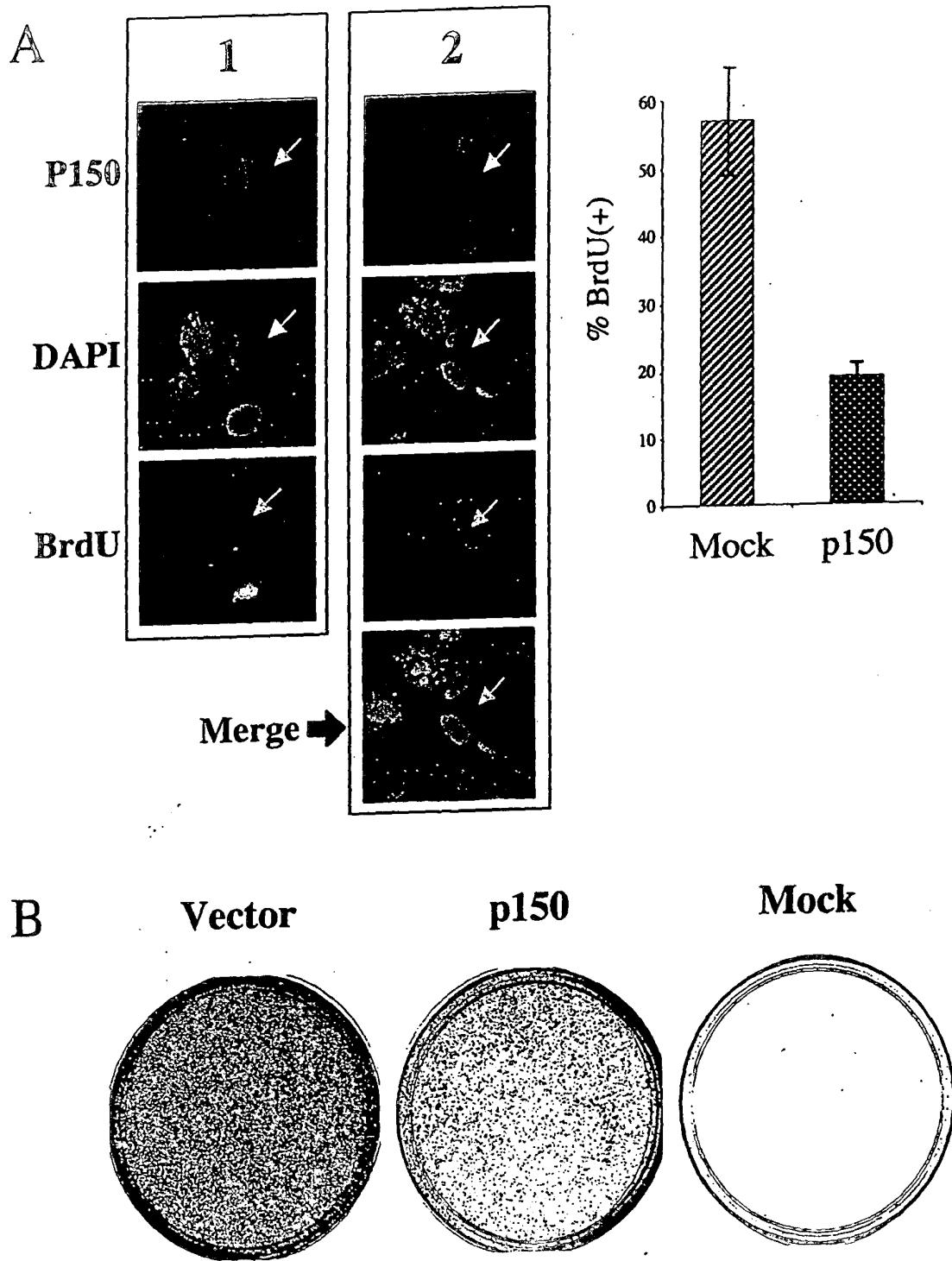


Fig. 10

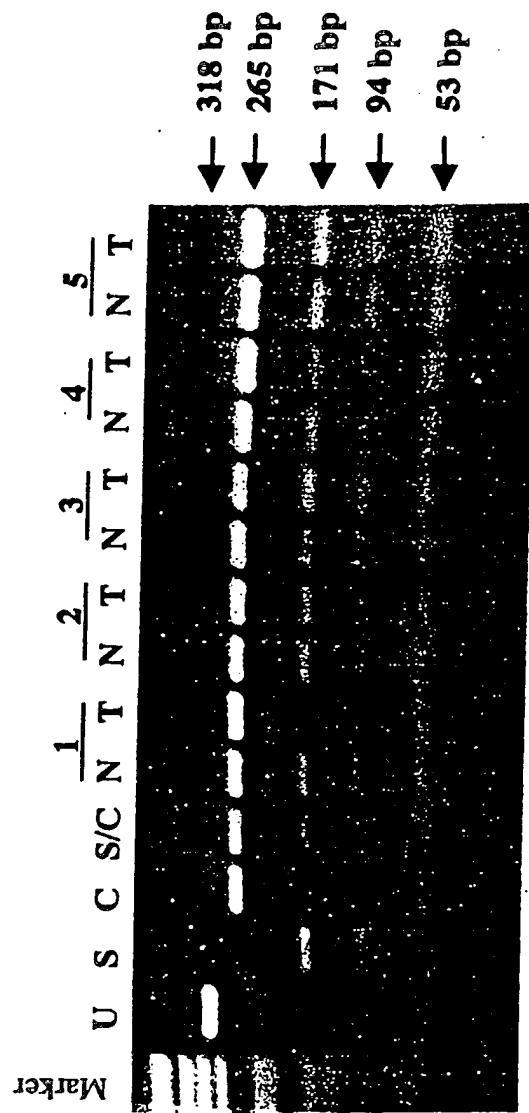


Fig. 11